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htgfer-II LTAETRITELGKOMLTAFHAKGALGEILTRIVISWEDLRNVGSSLARGLSHLHSDMTP-C
BACKR-IIB IAAETRISHLEVEILLITAFHDKGSLLDTIKGHITANELCHVAETHSRGISYLHEDVPMCR
BACKR-II IGAETRISVDVDIMLITAFHEKGSLSDFLARMVSMHELCHIAETHARGLAYLHEDIFGLK
dzi-1 IGSDXVDTGFVTELMLVIETHPSGSLADFLLENTVNIETYYNLARSTASGLAFLANQIGGSK
Subdomains

V VI-A

CONE.8A
DIX N DFG

HTGFBR-II -GRPROPIVIODIXSSHILVRODITCCLCDFGLSLRL---GPYSSVDDLANSGOVGTARYMAP

BACKR-IIB GEGHRPSIANDFFSRVVLLSDLTAVLADFGLAVF---EPGEPPGD--THGGVGTRRYMAP

BACKR-II -GRHPAISNDLTSRNVLLSDLTAVLADFGLAVF---BAGKSAGD--THGGVGTRRYMAP

BACKR-II - ESNEPAISNDLTSRNVLLSDLTAVLAGDLGLSLSKPEDAASDIIAN--ENYKCGTVRYLAP

BUDGGBAIDS VI-B VII VIII

## (57) Abstract

A new receptor family has been identified, of activin-like kinases. Novel proteins have activin/TGF-\(\beta\)-type I receptor functionality, and have consequential diagnostic/therapeutic utility. They may have a serine/threonine kinase domain, a DFKSRN or DLKSKN sequence in subdomain VIB and/or a GTKRYM sequence in subdomain VIII.